

SEQUENCE LISTING

<110> UAB Research Foundation

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WU, Hongju

<120> METHODS AND COMPOSITIONS FOR IN VIVO
INFLAMMATION MONITORING

<130> 21085.0050P1

<140> Unassigned
<141> 2004-09-23

<150> 60/505,543
<151> 2003-09-23

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<170> FastSEQ for Windows Version 4.0

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ccgttcccac	gccccggc	acgtcacaaa	ctccaccccc	tcattatcat	attggcttca	36000
atccaaaata	aggtatatta	tttatgtat				36029

<210> 6
<211> 720
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<221> misc_feature
<222> 693, 709
<223> n = g, a, c or t(u)

<400> 6

ttcaactagg	tgtcctcgga	tcaccacgaag	tgaaaaattaa	acacttttct	ccgtatcacg	60
aagtaaaaat	taaacacttt	tctccgtatg	gatcccatca	ccatcaccat	cacctagg	120
cacctaaata	tgccgataaa	acatttcaac	ctgaacctca	aataggagaa	tctcgttgt	180
acgaaacaga	aattaatcat	gcagctggga	gagtctaaa	aaagactacc	ccaatgaaac	240
catgttacgg	ttcatatgca	aaacccacaa	atgaaaaatgg	agggcaaggc	attcttgtaa	300
agcaacaaaa	tggaaagcta	gaaagtcaag	tggaaatgca	attttctca	actactgagg	360
cagccgcagg	caatggtgat	aacttgcactc	ctaaagtgg	attgtacagt	gaagatgtag	420
atataaaaaac	cccagacact	catatttctt	acatgccac	tattaaggaa	ggtaactcac	480
gagaactaat	gggocaacaa	totatgccc	acaggcctaa	ttacattgt	tttagggaca	540
attttattgg	tctaatgtat	tacaacagca	cggtaatat	gggtgttctg	gcgggccaag	600
catcgagtt	gaatgtgtt	gtagatttgc	aagacagaaa	cacagagctt	tcataaccagc	660
ttttgcttga	ttccattgg	gatagaacca	ggntacttt	ctatgtggna	tcaggcttgt	720

<210> 7
<211> 719
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<221> misc_feature
<222> 2,3,4,10,12,16,17,42,715,719
<223> n = g, a, c or t(u)

<400> 7

cnngggaggn cntcnata ggtgtcgaag gtcaaaccacc tnaatatgcc gataaaacat	60
ttcaacctga acctcaaata ggagaatctc agtgg tacga aacagaaaatt aatcatgcag	120
ctgggagagt cctaaaaaag actaccccaa tgaaaccatg ttacgggtca tatgcaaaac	180
ccacaaatga aaatggaggg caaggcattc ttgtaaaagca acaaaatgga aagctagaaa	240
gtcaagtggta aatgcaattt ttctcaacta ctctcggttc ccacgaagtg aaaattaaac	300
actttctcc gtatcacgaa gtgaaaatta aacactttc tccgtatgga tcccatcacc	360
atcaccatca cctaggttca ttgactccta aagtggattt gtacagtgaa gatgttagata	420
tagaaacccc agacactcat atttcttaca tgcccactat taaggaaggt aactcacgag	480
aactaatggg ccaacaatct atgccccaca ggcctaatta cattgctttt aggacaaatt	540
ttattggct aatgtattac aacagcacgg gtaatatggg tggtctggcg ggccaagcat	600
cgcagttgaa tgctgttga gatttgcag acagaaaacac agagtttca taccagctt	660
tgcttatttca cattgggtat agaaccaggta actttcttat gtggaatcag gctgntgan	719

<210> 8
<211> 108
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 8
ctcgatccc acgaagtgaa aattaaacac ttttccgt atcacgaagt gaaaattaaa
cacttttctc cgtatggatc ccatcaccat caccatcacc tagttca

60
108

<210> 9
<211> 36
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 9
Leu Gly Ser His Glu Val Lys Ile Lys His Phe Ser Pro Tyr His Glu
1 5 10 15
Val Lys Ile Lys His Phe Ser Pro Tyr Gly Ser His His His His His
20 25 30

His Leu Gly Ser
35

<210> 10
<211> 11
<212> PRT
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 10
His Glu Val Lys Ile Lys His Phe Ser Pro Tyr
1 5 10

<210> 11
<211> 5
<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 11

Gly Gly Gly Gly Ser
1 5

<210> 12

<211> 12

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 12

Leu Gly Ser His His His His His His Leu Gly Ser
1 5 10

<210> 13

<211> 3

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 13

Lys Gly Ser
1

<210> 14

<211> 22

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 14

cctacgcacg acgtgaccac ag

22

<210> 15

<211> 62

<212> DNA

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence; note =
synthetic construct

<400> 15

tgaacctagg tcatggat ggtgatggga tccgaggaca cctatttcaa taccctcctt
tg

60

62

<210> 16
<211> 61
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 16
ctcgatccc atcaccatca ccatcaccta gttcaccta aatatgccga taaaacattt 60
c 61

<210> 17
<211> 23
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 17
ctaggagct ctgcagaacc atg 23

<210> 18
<211> 60
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 18
tgaacctagg tcatggat ggtgatggga tccgagttcg taccactgag atttccttat 60

<210> 19
<211> 60
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 19
ctcgatccc atcaccatca ccatcaccta gttcaactg aaattaatca tgtagctggg 60

<210> 20
<211> 61
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 20
tgaacctagg tcatggat ggtgatggga tccgagagta gttgagaaaa attgcatttc 60
c 61

<210> 21
<211> 60
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 21
ctcggatccc atcaccatca ccatcaccta ggttcattga ctcctaaagt ggtattgtac 60

<210> 22
<211> 61
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 22
tgaacctagg tcatggtgat ggtgatggga tccgagagtg ggcattgttaag aaatatgttgt 60
g 61

<210> 23
<211> 58
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 23
ctcggatccc atcaccatca ccatcaccta ggttcaaact cacgagaact aatgggcc 58

<210> 24
<211> 60
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 24
tgaacctagg tcatggtgat ggtgatggga tccgagaggt tttaccttgg taagagtctc 60

<210> 25
<211> 62
<212> DNA
<213> Artificial Sequence

<220>
<223> Description of Artificial Sequence; note =
synthetic construct

<400> 25
ctcggatccc atcaccatca ccatcaccta ggttcatggg aaaaagatgc tacagaattt 60

synthetic construct

<221> misc_feature
 <222> 1-282, 390-720
 <223> n = g, a, c or t(u)

<400> 29

nnnnnnnnnn	60							
nnnnnnnnnn	120							
nnnnnnnnnn	180							
nnnnnnnnnn	240							
nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnctcgatc	ccacgaagt	aaaattaaac		300
acttttctcc	gtatcacgaa	gtgaaaatta	aacactttc	tccgtatgga	tcccatcacc			360
atcaccatca	cctaggttca	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	nnnnnnnnnn	420
nnnnnnnnnn	480							
nnnnnnnnnn	540							
nnnnnnnnnn	600							
nnnnnnnnnn	660							
nnnnnnnnnn	720							